

## SEQUENCE LISTING

<110> Sheppard, Paul O.  
 Baidur, Nand  
 Bishop, Paul D.

<120> MAMMALIAN ADHESION PROTEASE PEPTIDES

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<213> Homo sapiens

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Ala Gly Arg Asp Ala Leu Cys Gly Lys Leu Gln Cys Gln Gly Gly Lys	
570 575 580	
ccc agc ctg ctc gca ccg cac atg gtg cca gtg gac tct acc gtt cac	1830
Pro Ser Leu Leu Ala Pro His Met Val Pro Val Asp Ser Thr Val His	
585 590 595	
cta gat ggc cag gaa gtg act tgt cgg gga gcc ttg gca ctc ccc agt	1878
Leu Asp Gly Gln Glu Val Thr Cys Arg Gly Ala Leu Ala Leu Pro Ser	
600 605 610	
gcc cag ctg gac ctg ctt ggc ctg ggc ctg gta gag cca ggc acc cag	1926
Ala Gln Leu Asp Leu Leu Gly Leu Gly Leu Val Glu Pro Gly Thr Gln	
615 620 625 630	
tgt gga cct aga atg gtg tgc cag agc agg cgc tgc agg aag aat gcc	1974
Cys Gly Pro Arg Met Val Cys Gln Ser Arg Arg Cys Arg Lys Asn Ala	
635 640 645	
ttc cag gag ctt cag cgc tgc ctg act gcc tgc cac agc cac ggg gtt	2022
Phe Gln Glu Leu Gln Arg Cys Leu Thr Ala Cys His Ser His Gly Val	
650 655 660	
tgc aat agc aac cat aac tgc cac tgt gct cca ggc tgg gct cca ccc	2070
Cys Asn Ser Asn His Asn Cys His Cys Ala Pro Gly Trp Ala Pro Pro	
665 670 675	
ttc tgt gac aag cca ggc ttt ggt ggc agc atg gac agt ggc cct gtg	2118
Phe Cys Asp Lys Pro Gly Phe Gly Gly Ser Met Asp Ser Gly Pro Val	
680 685 690	
cag gct gaa aac cat gac acc ttc ctg ctg gcc atg ctc ctc agc gtc	2166
Gln Ala Glu Asn His Asp Thr Phe Leu Leu Ala Met Leu Leu Ser Val	
695 700 705 710	



ctg ctg cct ctg ctc cca ggg gcc ggc ctg gcc tgg tgt tgc tac cga Leu Leu Pro Leu Leu Pro Gly Ala Gly Leu Ala Trp Cys Cys Tyr Arg 715 720 725	2214
ctc cca gga gcc cat ctg cag cga tgc agc tgg ggc tgc aga agg gac Leu Pro Gly Ala His Leu Gln Arg Cys Ser Trp Gly Cys Arg Arg Asp 730 735 740	2262
cct gcg tgc agt ggc ccc aaa gat ggc cca cac agg gac cac ccc ctg Pro Ala Cys Ser Gly Pro Lys Asp Gly Pro His Arg Asp His Pro Leu 745 750 755	2310
ggc ggc gtt cac ccc atg gag ttg ggc ccc aca gcc act gga cag ccc Gly Gly Val His Pro Met Glu Leu Gly Pro Thr Ala Thr Gly Gln Pro 760 765 770	2358
tgg ccc ctg gac cct gag aac tct cat gag ccc agc agc cac cct gag Trp Pro Leu Asp Pro Glu Asn Ser His Glu Pro Ser Ser His Pro Glu 775 780 785 790	2406
aag cct ctg cca gca gtc tcg cct gac ccc caa gat caa gtc cag atg Lys Pro Leu Pro Ala Val Ser Pro Asp Pro Gln Asp Gln Val Gln Met 795 800 805	2454
cca aga tcc tgc ctc tgg tgagaggtag ctccataaaat gaacagattt Pro Arg Ser Cys Leu Trp 810	2502
aaagacaggt ggccactgac agccactcca ggaacttgaa ctgcaggggc agagccagtg aatcaccgga cctccagcac ctgcaggcag cttggaagtt tcttccccga gtggagcttc gacccacca ctccaggaaac ccagagccac attagaagtt cctgagggct ggagaacact gctgggcaca ctctccagct caataaacca tcagtcccag aagcaaaggt cacacagccc ctgacctccc tcaccagtgg aggctgggta gtgctggcca tccaaaagg gctctgtcct gggagtctgg tgtgtctcct acatgcaatt tccacggacc cagctctgtg gagggcatga ctgctggcca gaagctagtg gtcttggggc cctatgggtc gactgagtcc acactcccct ggagcctggc tggcctctgc aaacaaacat aattttgggg accttccttc ctgtttcttc ccaccctgtc ttctccccta ggtggttcct gagccccac cccaatccc agtgctacac ctgaggttct ggagctcaga atctgacagc ctctcccca ttctgtgtgt gtcgggggga cagagggaaac catttaagaa aagataccaa agtagaagtc aaaagaaaga catgttggt ataggcgtgg tggctcatgc ctataatccc agcactttgg gaagccgggg taggaggatc accagaggcc agcaggtcca caccagcctg ggcaacacag caagacaccg catctacaga aaaattttaa aattagctgg gcgtgggtgt gtgtacctgt aggcctagct gctcaggagg ctgaagcagg aggatcactt gagcctgagt tcaacactgc agtgagctat ggtggcacca	2562 2622 2682 2742 2802 2862 2922 2982 3042 3102 3162 3222 3282 3342 3402



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atatta 3468

<210> 4

<211> 812

<212> PRT

<213> Homo sapiens

<400> 4

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Leu	Leu	Leu	Leu	Leu	Leu	Trp	Pro	Val	Pro	Gly	Ala	Gly	Val	Leu	Gln	
			20				25					30				
Gly	His	Ile	Pro	Gly	Gln	Pro	Val	Thr	Pro	His	Trp	Val	Leu	Asp	Gly	
		35				40					45					
Gln	Pro	Trp	Arg	Thr	Val	Ser	Leu	Glu	Glu	Pro	Val	Ser	Lys	Pro	Asp	
	50				55			60								
Met	Gly	Leu	Val	Ala	Leu	Glu	Ala	Glu	Gly	Gln	Glu	Leu	Leu	Leu	Glu	
65				70				75						80		
Leu	Glu	Lys	Asn	His	Arg	Leu	Leu	Ala	Pro	Gly	Tyr	Ile	Glu	Thr	His	
			85					90					95			
Tyr	Gly	Pro	Asp	Gly	Gln	Pro	Val	Val	Leu	Ala	Pro	Asn	His	Thr	Asp	
		100					105					110				
His	Cys	His	Tyr	Gln	Gly	Arg	Val	Arg	Gly	Phe	Pro	Asp	Ser	Trp	Val	
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Val	Leu	Cys	Thr	Cys	Ser	Gly	Met	Ser	Gly	Leu	Ile	Thr	Leu	Ser	Arg	
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Asn	Ala	Ser	Tyr	Tyr	Leu	Arg	Pro	Trp	Pro	Pro	Arg	Gly	Ser	Lys	Asp	
145					150				155					160		
Phe	Ser	Thr	His	Glu	Ile	Phe	Arg	Met	Glu	Gln	Leu	Leu	Thr	Trp	Lys	
			165					170						175		
Gly	Thr	Cys	Gly	His	Arg	Asp	Pro	Gly	Asn	Lys	Ala	Gly	Met	Thr	Ser	
		180						185				190				
Leu	Pro	Gly	Gly	Pro	Gln	Ser	Arg	Gly	Arg	Arg	Glu	Ala	Arg	Arg	Thr	
	195						200					205				
Arg	Lys	Tyr	Leu	Glu	Leu	Tyr	Ile	Val	Ala	Asp	His	Thr	Leu	Phe	Leu	
	210					215					220					
Thr	Arg	His	Arg	Asn	Leu	Asn	His	Thr	Lys	Gln	Arg	Leu	Leu	Glu	Val	
225				230						235				240		
Ala	Asn	Tyr	Val	Asp	Gln	Leu	Leu	Arg	Thr	Leu	Asp	Ile	Gln	Val	Ala	
			245						250				255			
Leu	Thr	Gly	Leu	Glu	Val	Trp	Thr	Glu	Arg	Asp	Arg	Ser	Arg	Val	Thr	
		260						265					270			



Gln	Asp	Ala	Asn	Ala	Thr	Leu	Trp	Ala	Phe	Leu	Gln	Trp	Arg	Arg	Gly
	275						280					285			
Leu	Trp	Ala	Gln	Arg	Pro	His	Asp	Ser	Ala	Gln	Leu	Leu	Thr	Gly	Arg
	290					295					300				
Ala	Phe	Gln	Gly	Ala	Thr	Val	Gly	Leu	Ala	Pro	Val	Glu	Gly	Met	Cys
305					310					315					320
Arg	Ala	Glu	Ser	Ser	Gly	Gly	Val	Ser	Thr	Asp	His	Ser	Glu	Leu	Pro
				325					330					335	
Ile	Gly	Ala	Ala	Ala	Thr	Met	Ala	His	Glu	Ile	Gly	His	Ser	Leu	Gly
		340						345					350		
Leu	Ser	His	Asp	Pro	Asp	Gly	Cys	Cys	Val	Glu	Ala	Ala	Ala	Glu	Ser
		355					360					365			
Gly	Gly	Cys	Val	Met	Ala	Ala	Ala	Thr	Gly	His	Pro	Phe	Pro	Arg	Val
	370					375					380				
Phe	Ser	Ala	Cys	Ser	Arg	Arg	Gln	Leu	Arg	Ala	Phe	Phe	Arg	Lys	Gly
385					390					395					400
Gly	Gly	Ala	Cys	Leu	Ser	Asn	Ala	Pro	Asp	Pro	Gly	Leu	Pro	Val	Pro
				405					410					415	
Pro	Ala	Leu	Cys	Gly	Asn	Gly	Phe	Val	Glu	Ala	Gly	Glu	Glu	Cys	Asp
			420					425					430		
Cys	Gly	Pro	Gly	Gln	Glu	Cys	Arg	Asp	Leu	Cys	Cys	Phe	Ala	His	Asn
		435					440					445			
Cys	Ser	Leu	Arg	Pro	Gly	Ala	Gln	Cys	Ala	His	Gly	Asp	Cys	Cys	Val
		450				455					460				
Arg	Cys	Leu	Leu	Lys	Pro	Ala	Gly	Ala	Leu	Cys	Arg	Gln	Ala	Met	Gly
465					470					475					480
Asp	Cys	Asp	Leu	Pro	Glu	Phe	Cys	Thr	Gly	Thr	Ser	Ser	His	Cys	Pro
				485					490					495	
Pro	Asp	Val	Tyr	Leu	Leu	Asp	Gly	Ser	Pro	Cys	Ala	Arg	Gly	Ser	Gly
			500					505					510		
Tyr	Cys	Trp	Asp	Gly	Ala	Cys	Pro	Thr	Leu	Glu	Gln	Gln	Cys	Gln	Gln
		515					520					525			
Leu	Trp	Gly	Pro	Gly	Ser	His	Pro	Ala	Pro	Glu	Ala	Cys	Phe	Gln	Val
	530					535					540				
Val	Asn	Ser	Ala	Gly	Asp	Ala	His	Gly	Asn	Cys	Gly	Gln	Asp	Ser	Glu
545					550					555					560
Gly	His	Phe	Leu	Pro	Cys	Ala	Gly	Arg	Asp	Ala	Leu	Cys	Gly	Lys	Leu
				565				570						575	
Gln	Cys	Gln	Gly	Gly	Lys	Pro	Ser	Leu	Leu	Ala	Pro	His	Met	Val	Pro
			580					585					590		
Val	Asp	Ser	Thr	Val	His	Leu	Asp	Gly	Gln	Glu	Val	Thr	Cys	Arg	Gly
		595					600					605			



Ala Leu Ala Leu Pro Ser Ala Gln Leu Asp Leu Leu Gly Leu Gly Leu  
610 615 620  
Val Glu Pro Gly Thr Gln Cys Gly Pro Arg Met Val Cys Gln Ser Arg  
625 630 635 640  
Arg Cys Arg Lys Asn Ala Phe Gln Glu Leu Gln Arg Cys Leu Thr Ala  
645 650 655  
Cys His Ser His Gly Val Cys Asn Ser Asn His Asn Cys His Cys Ala  
660 665 670  
Pro Gly Trp Ala Pro Pro Phe Cys Asp Lys Pro Gly Phe Gly Gly Ser  
675 680 685  
Met Asp Ser Gly Pro Val Gln Ala Glu Asn His Asp Thr Phe Leu Leu  
690 695 700  
Ala Met Leu Leu Ser Val Leu Leu Pro Leu Leu Pro Gly Ala Gly Leu  
705 710 715 720  
Ala Trp Cys Cys Tyr Arg Leu Pro Gly Ala His Leu Gln Arg Cys Ser  
725 730 735  
Trp Gly Cys Arg Arg Asp Pro Ala Cys Ser Gly Pro Lys Asp Gly Pro  
740 745 750  
His Arg Asp His Pro Leu Gly Gly Val His Pro Met Glu Leu Gly Pro  
755 760 765  
Thr Ala Thr Gly Gln Pro Trp Pro Leu Asp Pro Glu Asn Ser His Glu  
770 775 780  
Pro Ser Ser His Pro Glu Lys Pro Leu Pro Ala Val Ser Pro Asp Pro  
785 790 795 800  
Gln Asp Gln Val Gln Met Pro Arg Ser Cys Leu Trp  
805 810

&lt;210&gt; 5

&lt;211&gt; 2406

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Degenerate sequence

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(2406)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 5

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ytnytnytny cngtnccngg ngcnngngtn ytnccargnc ayathccngg ncarccngtn 120  
acnccncayt gggtnytnga yggncarccn tggmgnacng tnwsnytnga rgarcngtn 180



wsnaarccng	ayatgggnyt	ngtngcnytn	gargcngarg	gncargaryt	nytnytngar	240
ytngaraara	aycaymgyt	nytnngcnccn	ggntayathg	aracncayta	yggncngay	300
ggncarccng	tngtnytngc	nccnaaycay	acngaycayt	gycaytayca	rggnmgngtn	360
mgnggnttyc	cngaywsntg	ggtngtynytn	tgyacntgyw	snggnatgws	nggnytnath	420
acnytnwsnm	gnaaygcnws	ntaytayytn	mgnccntggc	cncnmgngg	nwsnaargay	480
ttywsnacnc	aygarathtt	ymgnatggar	carytnytna	cntggaargg	nacntgyggg	540
caymgngayc	cnggnaayaa	rgcnggnatg	acnwsnytn	cnggnggncc	ncarwsnmgn	600
ggnmgnmgng	argcnmgng	nacnmgnaar	tayytngary	tntayathgt	ngcngaycay	660
acnytnntty	tnacnmgnca	ymgnaayytn	aaycayacna	arcarmgnyt	nytnngargtn	720
gcnaaytayg	tngaycaryt	nytnmgnaacn	ytngayathc	argtngcnyt	nacnggnytn	780
gargtntgga	cngarmgnga	ymgnwsnmgn	gtnacncarg	aygcnaaygc	nacnytntg	840
gcnttyytnc	artggmgnmg	nggnytntg	gcncarmgnc	cncaygayws	ngcncarytn	900
ytnacnggm	gngcnttyca	rgngcgnacn	gtnggnytn	cncngtnga	rggnatgtgy	960
mgngcngarw	snwsnggngg	ngtnwsnacn	gaycaywsng	arytnccnat	hggngcngcn	1020
gcnacnatgg	cncaygarat	hggncaywsn	ytnggnytnw	sncaygaycc	ngayggntgy	1080
tgygtngarg	cngcngcnga	rsnggnggn	tgygtnatgg	cngcngcnac	nggncayccn	1140
ttyccnmgng	tnttywsngc	ntgywsnmgn	mgncarytnm	gngcnttytt	ymgnaarggn	1200
ggnggngcnt	gyttnwsnaa	ygcncngay	ccnggnytn	cngtnccncc	ngcnytntg	1260
ggnaayggnt	tygtngargc	nggngargar	tgygaytgyg	gncnggnca	rgartgymgn	1320
gaytntgyt	gyttygcnc	yaaytgywsn	ytngmncng	gngcncartg	ygcncaygg	1380
gaytgytgyg	tnmgntgyyt	nytnaarccn	gcnggngcny	tntgymgnca	rgcnatgggn	1440
gaytgygayy	tnccngartt	ytgyacnggn	acnwsnwsnc	aytgyccncc	ngaygtntay	1500
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acnytn Garc	arcartgyca	rcarytntg	ggncnggnw	sncayccngc	nccngargcn	1620
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wsngaygcng	cnggngcngc	ngarggnacn	ytngmngcng	tngcncnca	ratggcncay	2220
acnggnacna	cncntgggc	ngcnttyacn	ccntggwsnt	gggncncna	rcnytnngay	2280
wsnccnggnc	cntggacny	nmgnacnytn	atgwsnccng	cngcnacny	nmgnwsnytn	2340
tgycarcarw	snmgnytnac	nccnaarath	aarwsnmgt	gycargaycc	ngcnwsnggn	2400
garmgn						2406

&lt;210&gt; 6

&lt;211&gt; 2439

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence



&lt;220&gt;

&lt;223&gt; Degenerate sequence

&lt;221&gt; misc\_feature

&lt;222&gt; (1)...(2439)

&lt;223&gt; n = A,T,C or G

&lt;400&gt; 6

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ytnytntgcc	cngtnccngg	ngcnggngtn	ytncarggnc	ayathccngg	ncarccngtn	120
acnccncayt	gggtnytnga	yggncarccn	tggmgnacng	tnwsnytnga	rgarccngtn	180
wsnaarccng	ayatgggnyt	ngtngcnytn	gargcngarg	gncargaryt	nytnytngar	240
ytngaraara	aycaymgyt	nytngcncn	ggntayathg	aracncayta	yggncngay	300
ggncarccng	tngtnytngc	nccnaaycay	acngaycayt	gycaytayca	rggnmgngtn	360
mgnggnttyc	cngaywsntg	ggtngtnytn	tgyacntgyw	snggnatgws	nggnytnath	420
acnytnwsnm	gnaaygcwsw	ntaytayytn	mgncntggc	cncnmgngg	nwsnaargay	480
ttywsnacnc	aygarathtt	ymgnatggar	carytnytna	cntggaargg	nacntgyggg	540
caymgngayc	cnggnaayaa	rgcnggnatg	acnwsnytn	cnggnggnc	ncarwsnmgn	600
ggnmgmngng	argcnmgmng	nacnmgnaar	tayytngary	tntayathgt	ngcngaycay	660
acnytnntty	tnacnmgnca	ymgnaayytn	aaycayacna	arcarmgnyt	nytnngargtn	720
gcnaaytayg	tngaycaryt	nytnmgncn	ytngayathc	argtngcnyt	nacnggnytn	780
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gcnttyytnc	artggmngm	nggnytntg	gcncarmgnc	cncaygayws	ngcncarytn	900
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mgngcngarw	snwsnggng	ngtnwsnacn	gaycaywsng	arytnccnat	hggngcngcn	1020
gcnacnatgg	cncaygarat	hggncaywsn	ytnggnytnw	sncaygaycc	ngayggntgy	1080
tgygtngarg	cngcngcnga	rwsnggnggn	tgygtnatgg	cngcngcnac	nggncayccn	1140
ttyccnmng	tnttywsngc	ntgywsnmgn	mgncarytnm	gngcnttytt	ymgnaarggn	1200
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mgntgymgna	araaygcntt	ycargarytn	carmgntgyy	tnacngcntg	ycaysncay	1980
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gcntggtgyt gytaymgyt nccnggngcn cayytnarm gntgywsntg gggntgymgn	2220
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aaywsncayg arccnwsnws ncayccngar aarcnytn cngcngtnws nccngayccn	2400
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&lt;210&gt; 7

&lt;211&gt; 23

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; oligonucleotide ZC17,993

&lt;400&gt; 7

gaagctcctg gaaggcattc ttc

23

&lt;210&gt; 8

&lt;211&gt; 21

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Oligonucleotide ZC17,994

&lt;400&gt; 8

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21

&lt;210&gt; 9

&lt;211&gt; 27

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Oligonucleotide API

&lt;400&gt; 9

ccatcctaatac gactcact atagggc

27

&lt;210&gt; 10

&lt;211&gt; 23



<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide APII

<400> 10

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23

<210> 11

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide ZC18,262

<400> 11

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<210> 12

<211> 18

<212> DNA

<213> Artificial Sequence

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<223> Oligonucleotide ZC22,481

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<210> 13

<211> 18

<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide ZC22,482

<400> 13

cgccatccca gcagtagc

18



<210> 14  
<211> 4  
<212> PRT  
<213> Homo sapiens

<400> 14  
Met Ser Glu Cys  
1

<210> 15  
<211> 4  
<212> PRT  
<213> Homo sapiens

<400> 15  
Arg Ser Glu Cys  
1

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